RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number: 09/734,583B
Source: 1FW16
Date Processed by STIC: 12/27/04

ENTERED

CRF Errors Edited by the STIC Systems Branch

Serial Number: 09/734,583B

CRF Edit Date: 12/29/04
Edited by: AR

___ Realigned nucleic acid/amino acid numbers/text in cases where the sequence text "wrapped" to the next line

___ Corrected the SEQ ID NO. Sequence numbers edited were:

_____

___ Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited:

_____

___ Deleted: ___ invalid beginning/end-of-file text ; ___ page numbers

___ Inserted mandatory headings/numeric identifiers, specifically:

_____

___ Moved responses to same line as heading/numeric identifier, specifically:

_____

✓ Other:
Seq. 8- changed <211> response to "12"
Seq. 2- added "1" after "positions" in <223> response
Seq. 10- inserted <220>, and <223> after 4th line in <223> response
globally corrected spelling of "bridge" and "residue"

Revised 09/09/2003

IFW16

RAW SEQUENCE LISTING DATE: 12/28/2004
PATENT APPLICATION: **US/09/734,583B** TIME: 16:29:42

Input Set : **A:\PTO.AMC.txt**
Output Set: **N:\CRF4\12282004\I734583B.raw**

```
      3 <110> APPLICANT: Hornik, Vered
      5 <120> TITLE OF INVENTION: CONFORMATIONALLY CONSTRAINED BACKBONE CYCLIZED SOMATOSTATIN
ANALOGS
      7 <130> FILE REFERENCE: 87534-3000
      9 <140> CURRENT APPLICATION NUMBER: 09/734,583B
     11 <141> CURRENT FILING DATE: 2000-12-13
     13 <160> NUMBER OF SEQ ID NOS: 10
     15 <170> SOFTWARE: PatentIn version 3.1
     17 <210> SEQ ID NO: 1
     18 <211> LENGTH: 14
     19 <212> TYPE: PRT
     20 <213> ORGANISM: mammalian
     22 <400> SEQUENCE: 1
     24 Ala Gly Cys Lys Asn Phe Phe Trp Lys Thr Phe Thr Ser Cys
     25 1               5                   10
     28 <210> SEQ ID NO: 2
     29 <211> LENGTH: 6
     30 <212> TYPE: PRT
     31 <213> ORGANISM: Artificial sequence
     33 <220> FEATURE:
W--> 34 <221> NAME/KEY: DISULFIDE BRIDGE
     35 <222> LOCATION: (1)..(1)
     36 <223> OTHER INFORMATION: Cys residues at amino acid positions 1 and 6 form a disulfide
bridge
     38 <220> FEATURE:
     39 <221> NAME/KEY: MOD_RES
     40 <222> LOCATION: (3)..(3)
     41 <223> OTHER INFORMATION: The Trp residue is the D isomer
     43 <220> FEATURE:
     44 <223> OTHER INFORMATION: Synthetic peptide
     46 <400> SEQUENCE: 2
     48 Cys Phe Trp Lys Thr Cys
     49 1               5
     52 <210> SEQ ID NO: 3
     53 <211> LENGTH: 6
     54 <212> TYPE: PRT
C--> 55 <213> ORGANISM: Artificial peptide
     57 <220> FEATURE:
     58 <221> NAME/KEY: MOD_RES
     59 <222> LOCATION: (1)..(1)
     60 <223> OTHER INFORMATION: N-Methyl
     62 <220> FEATURE:
     63 <221> NAME/KEY: MOD_RES
     64 <222> LOCATION: (1)..(6)
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65 <223> OTHER INFORMATION: cyclo

RAW SEQUENCE LISTING DATE: 12/28/2004
PATENT APPLICATION: **US/09/734,583B** TIME: 16:29:42

Input Set : **A:\PTO.AMC.txt**
Output Set: **N:\CRF4\12282004\I734583B.raw**

```
     67 <220> FEATURE:
     68 <221> NAME/KEY: MOD_RES
     69 <222> LOCATION: (3)..(3)
     70 <223> OTHER INFORMATION: The Trp residue is the D isomer
     72 <220> FEATURE:
     73 <223> OTHER INFORMATION: Synthetic peptide
     76 <400> SEQUENCE: 3
     78 Ala Tyr Trp Lys Val Phe
     79 1               5
     82 <210> SEQ ID NO: 4
     83 <211> LENGTH: 8
     84 <212> TYPE: PRT
     85 <213> ORGANISM: Artificial Sequence
     87 <220> FEATURE:
     88 <221> NAME/KEY: MOD_RES
     89 <222> LOCATION: (1)..(1)
     90 <223> OTHER INFORMATION: The Phe residue is a D isomer
     92 <220> FEATURE:
     93 <221> NAME/KEY: MOD_RES
     94 <222> LOCATION: (8)..(8)
     95 <223> OTHER INFORMATION: The Thr residue ends with CH2OH
     97 <220> FEATURE:
W--> 98 <221> NAME/KEY: DISULFIDE BRIDGE
     99 <222> LOCATION: (2)..(2)
    100 <223> OTHER INFORMATION: A disulfide bridge is formed between Cys residues 2 and 7
    102 <220> FEATURE:
    103 <221> NAME/KEY: MOD_RES
    104 <222> LOCATION: (4)..(4)
    105 <223> OTHER INFORMATION: The Trp residue is a D isomer
    107 <220> FEATURE:
    108 <223> OTHER INFORMATION: Synthetic peptide
    111 <400> SEQUENCE: 4
    113 Phe Cys Phe Trp Lys Thr Cys Thr
    114 1               5
    117 <210> SEQ ID NO: 5
    118 <211> LENGTH: 7
    119 <212> TYPE: PRT
    120 <213> ORGANISM: Artificial Sequence
    122 <220> FEATURE:
W--> 123 <221> NAME/KEY: DISULFIDE
    124 <222> LOCATION: (2)..(2)
    125 <223> OTHER INFORMATION: A Disulfide Bridge is formed between the Cys residues at
position
    126       2 and 6
    128 <220> FEATURE:
    129 <221> NAME/KEY: MOD_RES
    130 <222> LOCATION: (1)..(1)
    131 <223> OTHER INFORMATION: The Phe residue is a D isomer
    133 <220> FEATURE:
    134 <221> NAME/KEY: MOD_RES
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RAW SEQUENCE LISTING DATE: 12/28/2004
PATENT APPLICATION: **US/09/734,583B** TIME: 16:29:42

Input Set : **A:\PTO.AMC.txt**
Output Set: **N:\CRF4\12282004\I734583B.raw**

```
135 <222> LOCATION: (4)..(4)
136 <223> OTHER INFORMATION: The Trp residue is a D isomer
138 <220> FEATURE:
139 <221> NAME/KEY: MOD_RES
140 <222> LOCATION: (7)..(7)
141 <223> OTHER INFORMATION: The Thr residue ends with N2H
143 <220> FEATURE:
144 <223> OTHER INFORMATION: Synthetic peptide
147 <400> SEQUENCE: 5
149 Phe Cys Phe Trp Lys Cys Thr
150 1               5
153 <210> SEQ ID NO: 6
154 <211> LENGTH: 8
155 <212> TYPE: PRT
156 <213> ORGANISM: Artificial sequence
158 <220> FEATURE:
159 <221> NAME/KEY: MISC_FEATURE
160 <222> LOCATION: (1)..(1)
161 <223> OTHER INFORMATION: is a gamma amino butyric acid, diamino butyric acid, Gly,
beta-Al
162       a, 5-amino pentanoic acid or amino hexanoic acid; Residue 1 is
163       bridged to Residue 8; Residue 1 also begins with a hydrogen, or a
164       mono- or di- saccharide attached
166 <220> FEATURE:
167 <221> NAME/KEY: MISC_FEATURE
168 <222> LOCATION: (2)..(2)
169 <223> OTHER INFORMATION: is (D) or (L) Phe or Tyr
171 <220> FEATURE:
172 <221> NAME/KEY: MISC_FEATURE
173 <222> LOCATION: (3)..(3)
174 <223> OTHER INFORMATION: is (D) or (L)-Trp, or (L)-Phe, (D)- or (L)-1Nal or (D) or
175       (L)-2Nal, or Tyr
177 <220> FEATURE:
178 <221> NAME/KEY: MISC_FEATURE
179 <222> LOCATION: (4)..(4)
180 <223> OTHER INFORMATION: is (D) or (L)-Trp
182 <220> FEATURE:
183 <221> NAME/KEY: MISC_FEATURE
184 <222> LOCATION: (5)..(5)
185 <223> OTHER INFORMATION: is (D) or (L)-Lys
187 <220> FEATURE:
188 <221> NAME/KEY: MISC_FEATURE
189 <222> LOCATION: (6)..(6)
190 <223> OTHER INFORMATION: is Thr, Gly, Abu, Ser, Cys, Val, (D) or (L)-Ala, or (D)- or
191       (L)-Ala, or Tyr
193 <220> FEATURE:
194 <221> NAME/KEY: MISC_FEATURE
195 <222> LOCATION: (7)..(7)
196 <223> OTHER INFORMATION: is (D) or (L)-Phe, or (D)- or (L)-Ala, Nle, or Cys
198 <220> FEATURE:
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RAW SEQUENCE LISTING DATE: 12/28/2004
PATENT APPLICATION: **US/09/734,583B** TIME: 16:29:42

Input Set : **A:\PTO.AMC.txt**
Output Set: **N:\CRF4\12282004\I734583B.raw**

```
    199 <221> NAME/KEY: MISC_FEATURE
    200 <222> LOCATION: (8)..(8)
    201 <223> OTHER INFORMATION: is Gly, Val, Leu, (D) or (L)-Phe, or 1Nal or 2Nal; with a
    202       terminal carboxy acid, amide or alcohol group
    204 <220> FEATURE:
    205 <223> OTHER INFORMATION: Synthetic peptide
    208 <400> SEQUENCE: 6
W--> 210 Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa
    211 1               5
    214 <210> SEQ ID NO: 7
    215 <211> LENGTH: 7
    216 <212> TYPE: PRT
    217 <213> ORGANISM: Artificial Sequence
    219 <220> FEATURE:
    220 <221> NAME/KEY: MISC_FEATURE
    221 <222> LOCATION: (1)..(1)
    222 <223> OTHER INFORMATION: is (D)- or (L)-Phe, or (D)- or (L)-Ala; wherein Residue 1 is
    223       bridged to Residue 6 a bridging group composed of 1 to 5 methyl
    224       spacers connected to an amide, thioether, thioester, or disulfide,
    225       followed by 1 to 5 methyl spacers
    227 <220> FEATURE:
    228 <221> NAME/KEY: MISC_FEATURE
    229 <222> LOCATION: (2)..(2)
    230 <223> OTHER INFORMATION: is Tyr or (D)- or (L)-Phe
    232 <220> FEATURE:
    233 <221> NAME/KEY: MISC_FEATURE
    234 <222> LOCATION: (3)..(3)
    235 <223> OTHER INFORMATION: is (D)- or (L)-Trp, (D)- or (L)-1Nal, or (D)- or (L)-2Nal
    237 <220> FEATURE:
    238 <221> NAME/KEY: MISC_FEATURE
    239 <222> LOCATION: (5)..(5)
    240 <223> OTHER INFORMATION: is Thr, Val, Ser, or Cys
    242 <220> FEATURE:
    243 <221> NAME/KEY: MISC_FEATURE
    244 <222> LOCATION: (6)..(6)
    245 <223> OTHER INFORMATION: is Gly or (D)- or (L)-Phe
    247 <220> FEATURE:
    248 <221> NAME/KEY: MISC_FEATURE
    249 <222> LOCATION: (7)..(7)
    250 <223> OTHER INFORMATION: is Thr, GABA, (D)- or (L)-1Nal, (D)- or (L)-2Nal, or (D)- or
    251       (L)-Phe
    253 <220> FEATURE:
    254 <223> OTHER INFORMATION: Synthetic peptide
    257 <400> SEQUENCE: 7
W--> 259 Xaa Xaa Xaa Lys Xaa Xaa Xaa
    260 1               5
    263 <210> SEQ ID NO: 8
    264 <211> LENGTH: 12
    265 <212> TYPE: PRT
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RAW SEQUENCE LISTING DATE: 12/28/2004
PATENT APPLICATION: **US/09/734,583B** TIME: 16:29:42

Input Set : **A:\PTO.AMC.txt**
Output Set: **N:\CRF4\12282004\I734583B.raw**

```
266 <213> ORGANISM: Artificial Sequence
268 <220> FEATURE:
269 <221> NAME/KEY: MISC_FEATURE
270 <222> LOCATION: (1)..(1)
271 <223> OTHER INFORMATION: is absent or is any amino acid
273 <220> FEATURE:
274 <221> NAME/KEY: MISC_FEATURE
275 <222> LOCATION: (2)..(2)
276 <223> OTHER INFORMATION: is absent or is any amino acid
278 <220> FEATURE:
279 <221> NAME/KEY: MISC_FEATURE
280 <222> LOCATION: (3)..(3)
281 <223> OTHER INFORMATION: is absent or is any amino acid
283 <220> FEATURE:
284 <221> NAME/KEY: MISC_FEATURE
285 <222> LOCATION: (4)..(4)
286 <223> OTHER INFORMATION: is absent or is any amino acid
288 <220> FEATURE:
289 <221> NAME/KEY: MISC_FEATURE
290 <222> LOCATION: (5)..(5)
291 <223> OTHER INFORMATION: is 1Nal, 2Nal, Beta-Asp (Ind), Gly, Tyr, (D)- or (L)-Ala, or
292       (D)- or (L)-Phe
294 <220> FEATURE:
295 <221> NAME/KEY: MISC_FEATURE
296 <222> LOCATION: (6)..(7)
297 <223> OTHER INFORMATION: may be absent, or are independently Gly, Tyr, 1Nal, 2Nal,
298       Beta-Asp (Ind), Gly, Tyr, (D)- or (L)-Ala, or (D)- or (L)-Phe
300 <220> FEATURE:
301 <221> NAME/KEY: MISC_FEATURE
302 <222> LOCATION: (8)..(8)
303 <223> OTHER INFORMATION: (D)- or (L)-Trp
305 <220> FEATURE:
306 <221> NAME/KEY: MISC_FEATURE
307 <222> LOCATION: (9)..(9)
308 <223> OTHER INFORMATION: (D)- or (L)-Lys
310 <220> FEATURE:
311 <221> NAME/KEY: MISC_FEATURE
312 <222> LOCATION: (10)..(10)
313 <223> OTHER INFORMATION: is absent or is Gly, Abu, Cys, Thr, Val, (D)- or (L)-Ala, or
314       (D)- or (L)-Phe
316 <220> FEATURE:
317 <221> NAME/KEY: MISC_FEATURE
318 <222> LOCATION: (11)..(11)
319 <223> OTHER INFORMATION: is Cys, (D)- or (L)-Ala, or (D)- or (L)-Phe
321 <220> FEATURE:
322 <221> NAME/KEY: MISC_FEATURE
323 <222> LOCATION: (12)..(12)
324 <223> OTHER INFORMATION: is absent or is Val, Thr, 1Nal or 2Nal
326 <220> FEATURE:
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RAW SEQUENCE LISTING ERROR SUMMARY DATE: 12/28/2004
PATENT APPLICATION: **US/09/734,583B** TIME: 16:29:43

Input Set : **A:\PTO.AMC.txt**
Output Set: **N:\CRF4\12282004\I734583B.raw**

Please Note:
Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:6; Xaa Pos. 1,2,3,4,5,6,7,8
Seq#:7; Xaa Pos. 1,2,3,5,6,7
Seq#:8; Xaa Pos. 1,2,3,4,5,6,7,8,9,10,11,12
Seq#:9; Xaa Pos. 1,2,3,4,5,6,7
Seq#:10; Xaa Pos. 1,2,3,4,5,6,7

VERIFICATION SUMMARY DATE: 12/28/2004
PATENT APPLICATION: **US/09/734,583B** TIME: 16:29:43

Input Set : **A:\PTO.AMC.txt**
Output Set: **N:\CRF4\12282004\I734583B.raw**

L:34 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:2
L:55 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:3
L:98 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:4
L:123 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:5
L:210 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6 after pos.:0
L:259 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7 after pos.:0
L:332 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8 after pos.:0
L:384 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:0
L:400 M:283 W: Missing Blank Line separator, <220> field identifier
L:439 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10 after pos.:0

IFW16

RAW SEQUENCE LISTING DATE: 12/27/2004
PATENT APPLICATION: **US/09/734,583B** TIME: 12:16:46

Input Set : **A:\87534-3000.txt**
Output Set: **N:\CRF4\12272004\I734583B.raw**

3 <110> APPLICANT: Hornik, Vered
5 <120> TITLE OF INVENTION: CONFORMATIONALLY CONSTRAINED BACKBONE CYCLIZED SOMATOSTATIN ANALOGS
7 <130> FILE REFERENCE: 87534-3000
9 <140> CURRENT APPLICATION NUMBER: 09/734,583B
11 <141> CURRENT FILING DATE: 2000-12-13
13 <160> NUMBER OF SEQ ID NOS: 10
15 <170> SOFTWARE: PatentIn version 3.1

Does Not Comply
Corrected Diskette Needed

rr 1, 3-5

ERRORED SEQUENCES

263 <210> SEQ ID NO: 8
264 <211> LENGTH: (9) 12
265 <212> TYPE: PRT
266 <213> ORGANISM: Artificial Sequence
268 <220> FEATURE:
269 <221> NAME/KEY: MISC_FEATURE
270 <222> LOCATION: (1)..(1)
271 <223> OTHER INFORMATION: is absent or is any amino acid
273 <220> FEATURE:
274 <221> NAME/KEY: MISC_FEATURE
275 <222> LOCATION: (2)..(2)
276 <223> OTHER INFORMATION: is absent or is any amino acid
278 <220> FEATURE:
279 <221> NAME/KEY: MISC_FEATURE
280 <222> LOCATION: (3)..(3)
281 <223> OTHER INFORMATION: is absent or is any amino acid
283 <220> FEATURE:
284 <221> NAME/KEY: MISC_FEATURE
285 <222> LOCATION: (4)..(4)
286 <223> OTHER INFORMATION: is absent or is any amino acid
288 <220> FEATURE:
289 <221> NAME/KEY: MISC_FEATURE
290 <222> LOCATION: (5)..(5)
291 <223> OTHER INFORMATION: is 1Nal, 2Nal, Beta-Asp (Ind), Gly, Tyr, (D)- or (L)-Ala, or
292 (D)- or (L)-Phe
294 <220> FEATURE:
295 <221> NAME/KEY: MISC_FEATURE
296 <222> LOCATION: (6)..(7)
297 <223> OTHER INFORMATION: may be absent, or are independently Gly, Tyr, 1Nal, 2Nal,
298 Beta-Asp (Ind), Gly, Tyr, (D)- or (L)-Ala, or (D)- or (L)-Phe
300 <220> FEATURE:
301 <221> NAME/KEY: MISC_FEATURE

RAW SEQUENCE LISTING DATE: 12/27/2004
PATENT APPLICATION: **US/09/734,583B** TIME: 12:16:46

Input Set : **A:\87534-3000.txt**
Output Set: **N:\CRF4\12272004\I734583B.raw**

302 <222> LOCATION: (8)..(8)
303 <223> OTHER INFORMATION: (D)- or (L)-Trp
305 <220> FEATURE:
306 <221> NAME/KEY: MISC_FEATURE
307 <222> LOCATION: (9)..(9)
308 <223> OTHER INFORMATION: (D)- or (L)-Lys
310 <220> FEATURE:
311 <221> NAME/KEY: MISC_FEATURE
312 <222> LOCATION: (10)..(10)
313 <223> OTHER INFORMATION: is absent or is Gly, Abu, Cys, Thr, Val, (D)- or (L)-Ala, or
314 (D)- or (L)-Phe
316 <220> FEATURE:
317 <221> NAME/KEY: MISC_FEATURE
318 <222> LOCATION: (11)..(11)
319 <223> OTHER INFORMATION: is Cys, (D)- or (L)-Ala, or (D)- or (L)-Phe
321 <220> FEATURE:
322 <221> NAME/KEY: MISC_FEATURE
323 <222> LOCATION: (12)..(12)
324 <223> OTHER INFORMATION: is absent or is Val, Thr, 1Nal or 2Nal
326 <220> FEATURE:
327 <223> OTHER INFORMATION: Synthetic peptide
330 <400> SEQUENCE: 8
W--> 332 **Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa**
E--> 333 **1 5 10**

<210> 2
<211> 6
<212> PRT
<213> Artificial sequence

<220>
<221> DISULFIDE BRIDGE
<222> (1)..(1)
<223> Cys residues at amino acid positions and 6 form a disulfide bridge

insert "1"

<222> (1)..(1)
<223> (D)- or (L)-Phe, Tyr or (D)- or (L)-Ala; Residue 1 is connected to Residue 7 by a bridge comprised of 1 to 5 methylene spacers conncected to an amide, thioether, thioester, or disulfide, followed by 1 to 5 methylene spacers

<220>
<221> MISC_FEATURE
<222> (2)..(2)
<223> (D)- or (L)-Phe, Tyr or (D)- or (L)-Ala;

<220>
<221> MISC_FEATURE
<222> (3)..(3)
<223> is absent or is (D)- or (L)-Phe, Tyr or (D)- or (L)-Ala

<220>
<221> MISC_FEATURE
<222> (4)..(4)
<223> is (D)- or (L)Tyr

<220>
<221> MISC_FEATURE
<222> (5)..(5)
<223> is (D)- or (L)-Lys

<220>
<221> MISC_FEATURE
<222> (6)..(6)
<223> is absent or is Thr, Val, Cys or (D)- or (L)-Ala

<220>
<221> MISC_FEATURE
<222> (7)..(7)
<223> is a (D)- or (L)-Phe, Cys, or (D)- or (L)-Ala

<220>
<223> Synthetic peptide

<400> 9

Xaa Xaa Xaa Xaa Xaa Xaa Xaa
1 5

<210> 10
<211> 7
<212> PRT
<213> Artificial Sequence

<220>
<221> MISC_FEATURE
<222> (1)..(1)
<223> is absent or is (D)- or (L)-Phe or Ala; the bride is connected to Residue 1 or 2 and Residue 6 or 7, wherein the birdge is comprised of 1 to 5 methylene spacers connected to an amide, thioether, thioester, or disulfide, followed by 1 to 5 methylene

bridge
bridge

see p. 5

<2237 response has a maximum of 4 lines

09/734,583B

5

insert <2207

insert <2237 spacers

insert <2207 after 4th line and <2237 at beginning of 5th line

<220>
<221> MISC_FEATURE
<222> (2)..(2)
<223> is (D)- or (L)-Phe or Ala, Tyr

<220>
<221> MISC_FEATURE
<222> (3)..(3)
<223> is (D)- or (L)-Trp

<220>
<221> MISC_FEATURE
<222> (4)..(4)
<223> is (D)- or (L)-Lys

<220>
<221> MISC_FEATURE
<222> (5)..(5)
<223> is Thr, Ala, Val, or Cys

<220>
<221> MISC_FEATURE
<222> (6)..(6)
<223> is absent or is (D)- or (L)-Phe, Ala, or Cys

<220>
<221> MISC_FEATURE
<222> (7)..(7)
<223> is absent or is Thr or Thr reduced to an alcohol

<220>
<223> Synthetic peptide

<400> 10

Xaa Xaa Xaa Xaa Xaa Xaa Xaa
1 5

VERIFICATION SUMMARY DATE: 12/27/2004
PATENT APPLICATION: **US/09/734,583B** TIME: 12:16:47

Input Set : **A:\87534-3000.txt**
Output Set: **N:\CRF4\12272004\I734583B.raw**

L:34 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:2
L:55 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:3
L:98 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:4
L:123 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:5
L:210 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6 after pos.:0
L:259 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7 after pos.:0
L:332 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8 after pos.:0
L:333 M:252 E: No. of Seq. differs, <211> LENGTH:Input:9 Found:12 SEQ:8
L:384 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:0
L:400 M:259 W: Allowed number of lines exceeded, <223> Other Information:
L:438 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10 after pos.:0